SEQUENCE LISTING

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<212> PRT

Irani, Meher Andersson, Claes Weigelt, Cecilia Christer Möller Pia Hydén <120> PRODUCTION AND PURIFICATION OF ARYL SULFATASE A <130> 33686PC01 <150> PA200400144 <151> 2004-01-30 <150> US 60/540,061 <151> 2004-01-30 <160> 4 <170> FastSEQ for Windows Version 4.0 <210> 1 <211> 1524 <212> DNA <213> Homo Sapiens <400> 1 atgggggeac cgcggtccct cctcctggcc ctggctgctg gcctggccgt tgcacgtccg 60 cccaacatcg tgctgatctt tgccgacgac ctcggctatg gggacctggg ctgctatggg 120 cacccaget ctaccactcc caacctggac cagetggcgg cgggagggct gcggttcaca 180 gacttctacg tgcctgtgtc tctgtgcaca ccctctaggg ccgccctcct gaccggccgg 240 etcecqqttc qqatqqcat qtacectqqc qtectqqtqc ccaqeteccq qqqqqgectg 300 cccctggagg aggtgaccgt ggccgaagtc ctggctgccc gaggctacct cacaggaatg 360 gccggcaagt ggcaccttgg ggtggggcct gagggggcct tcctgccccc ccatcagggc 420 ttccatcgat ttctaggcat cccgtactcc cacgaccagg gcccctgcca gaacctgacc 480 tgcttcccgc cggccactcc ttgcgacggt ggctgtgacc agggcctggt ccccatccca 540 ctgttggcca acctgtccgt ggaggcgcag ccccctggc tgcccggact agaggcccgc 600 tacatggctt tegeccatga ceteatggce gaegeccage gecaggateg eccettette 660 ctgtactatg cctctcacca cacccactac cctcagttca gtgggcagag ctttgcagag 720 cqttcagqcc gcqqgccatt tggggactcc ctgatggagc tggatgcagc tgtggggacc 780 ctgatgacag ccatagggga cctggggctg cttgaagaga cgctggtcat cttcactgca 840 qacaatqqac ctqaqaccat gcgtatgtcc cgaggcggct gctccggtct cttgcggtgt 900 qqaaaqqqaa cqacctacqa qqqcqqtqtc cqaqaqcctq ccttqqcctt ctgqccaggt 960 catategete eeggegtgae eeacgagetg gecageteee tggacetget gectaceetg1020 qcaqccctqq ctqqqqccc actqcccaat qtcaccttqq atqqctttqa cctcaqcccc1080 ctgctgctgg gcacaggcaa gagcctcgg cagtctctct tcttctaccc gtcctaccca1140 gacgaggtcc gtggggtttt tgctgtgcgg actggaaagt acaaggctca cttcttcacc1200 cagggetetg eccaeagtga taccaetgea gaccetgeet gecaegeete cagetetetg1260 actgctcatg agccccgct gctctatgac ctgtccaagg accctggtga gaactacaac1320 ctgctggggg gtgtggccgg ggccacccca gaggtgctgc aagccctgaa acagcttcag1380 ctgctcaagg cccagttaga cgcagctgtg accttcggcc ccagccaggt ggcccggggc1440 qaqqaccccq ccctgcagat ctgctgtcat cctggctgca cccccgccc agcttgctgc1500 cattgcccag atccccatgc ctga <210> 2 <211> 507

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1 5
Asp Leu Gly Cys Tyr Gly His

Arg Pro Pro Asn Ile Val Leu Ile Phe Ala Asp Asp Leu Gly Tyr Gly 10 Asp Leu Gly Cys Tyr Gly His Pro Ser Ser Thr Thr Pro Asn Leu Asp 2.5 Gln Leu Ala Ala Gly Gly Leu Arg Phe Thr Asp Phe Tyr Val Pro Val 40 Ser Leu Xaa Thr Pro Ser Arg Ala Ala Leu Leu Thr Gly Arg Leu Pro Val Arg Met Gly Met Tyr Pro Gly Val Leu Val Pro Ser Ser Arg Gly 75 Gly Leu Pro Leu Glu Glu Val Thr Val Ala Glu Val Leu Ala Ala Arg 90 Gly Tyr Leu Thr Gly Met Ala Gly Lys Trp His Leu Gly Val Gly Pro 100 105 Glu Gly Ala Phe Leu Pro Pro His Gln Gly Phe His Arg Phe Leu Gly 120 Ile Pro Tyr Ser His Asp Gln Gly Pro Cys Gln Asn Leu Thr Cys Phe 135 Pro Pro Ala Thr Pro Cys Asp Gly Gly Cys Asp Gln Gly Leu Val Pro 145 150 155 Ile Pro Leu Leu Ala Asn Leu Ser Val Glu Ala Gln Pro Pro Trp Leu 165 170 Pro Gly Leu Glu Ala Arg Tyr Met Ala Phe Ala His Asp Leu Met Ala 180 185 Asp Ala Gln Arg Gln Asp Arg Pro Phe Phe Leu Tyr Tyr Ala Ser His 200 205 His Thr His Tyr Pro Gln Phe Ser Gly Gln Ser Phe Ala Glu Arg Ser 215 220 Gly Arg Gly Pro Phe Gly Asp Ser Leu Met Glu Leu Asp Ala Ala Val 230 235 Gly Thr Leu Met Thr Ala Ile Gly Asp Leu Gly Leu Leu Glu Glu Thr 250 245 Leu Val Ile Phe Thr Ala Asp Asn Gly Pro Glu Thr Met Arg Met Ser 265 Arg Gly Gly Cys Ser Gly Leu Leu Arg Cys Gly Lys Gly Thr Thr Tyr 280 Glu Gly Gly Val Arg Glu Pro Ala Leu Ala Phe Trp Pro Gly His Ile 295 300 Ala Pro Gly Val Thr His Glu Leu Ala Ser Ser Leu Asp Leu Leu Pro 310 315 Thr Leu Ala Ala Leu Ala Gly Ala Pro Leu Pro Asn Val Thr Leu Asp 325 330

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Gly Phe Asp Leu Ser Pro Leu Leu Gly Thr Gly Lys Ser Pro Arg 345 Gln Ser Leu Phe Phe Tyr Pro Ser Tyr Pro Asp Glu Val Arg Gly Val 360 Phe Ala Val Arg Thr Gly Lys Tyr Lys Ala His Phe Phe Thr Gln Gly 375 380 Ser Ala His Ser Asp Thr Thr Ala Asp Pro Ala Cys His Ala Ser Ser 390 395 Ser Leu Thr Ala His Glu Pro Pro Leu Leu Tyr Asp Leu Ser Lys Asp 405 410 Pro Gly Glu Asn Tyr Asn Leu Leu Gly Gly Val Ala Gly Ala Thr Pro 420 425 Glu Val Leu Gln Ala Leu Lys Gln Leu Gln Leu Leu Lys Ala Gln Leu 440 Asp Ala Ala Val Thr Phe Gly Pro Ser Gln Val Ala Arg Gly Glu Asp 455 460 Pro Ala Leu Gln Ile Cys Cys His Pro Gly Cys Thr Pro Arg Pro Ala 470 475 Cys Cys His Cys Pro Asp Pro His Ala 485

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<212> PRT

<213> Homo Sapiens

<400> 4

Arg Pro Pro Asn Ile Val Leu Ile Phe Ala Asp Asp Leu Gly Tyr Gly 10 Asp Leu Gly Cys Tyr Gly His Pro Ser Ser Thr Thr Pro Asn Leu Asp 2.5 Gln Leu Ala Ala Gly Gly Leu Arg Phe Thr Asp Phe Tyr Val Pro Val 40 4.5 Ser Leu Cys Thr Pro Ser Arg Ala Ala Leu Leu Thr Gly Arg Leu Pro 55 Val Arg Met Gly Met Tyr Pro Gly Val Leu Val Pro Ser Ser Arg Gly 70 75 Gly Leu Pro Leu Glu Glu Val Thr Val Ala Glu Val Leu Ala Arg 85 90 Gly Tyr Leu Thr Gly Met Ala Gly Lys Trp His Leu Gly Val Gly Pro 100 105 Glu Gly Ala Phe Leu Pro Pro His Gln Gly Phe His Arg Phe Leu Gly 120 Ile Pro Tyr Ser His Asp Gln Gly Pro Cys Gln Asn Leu Thr Cys Phe 135 Pro Pro Ala Thr Pro Cys Asp Gly Gly Cys Asp Gln Gly Leu Val Pro 150 155 Ile Pro Leu Leu Ala Asn Leu Ser Val Glu Ala Gln Pro Pro Trp Leu 165 170 Pro Gly Leu Glu Ala Arg Tyr Met Ala Phe Ala His Asp Leu Met Ala 180 185 Asp Ala Gln Arg Gln Asp Arg Pro Phe Phe Leu Tyr Tyr Ala Ser His 205 200 His Thr His Tyr Pro Gln Phe Ser Gly Gln Ser Phe Ala Glu Arg Ser 220 215 Gly Arg Gly Pro Phe Gly Asp Ser Leu Met Glu Leu Asp Ala Ala Val 230 235 Gly Thr Leu Met Thr Ala Ile Gly Asp Leu Gly Leu Leu Glu Glu Thr 245 250 Leu Val Ile Phe Thr Ala Asp Asn Gly Pro Glu Thr Met Arg Met Ser

			260					265					270		
Arg	Gly	Gly 275	Cys	Ser	Gly	Leu	Leu 280	Arg	Cys	Gly	Lys	Gly 285	Thr	Thr	Tyr
Glu	Gly 290	Gly	Val	Arg	Glu	Pro 295	Ala	Leu	Ala	Phe	Trp 300	Pro	Gly	His	Ile
Ala 305	Pro	Gly	Val	Thr	His 310	Glu	Leu	Ala	Ser	Ser 315	Leu	Asp	Leu	Leu	Pro 320
Thr	Leu	Ala	Ala	Leu 325	Ala	Gly	Ala	Pro	Leu 330	Pro	Asn	Val	Thr	Leu 335	Asp
Gly	Phe	Asp	Leu 340	Ser	Pro	Leu	Leu	Leu 345	Gly	Thr	Gly	Lys	Ser 350	Pro	Arg
Gln	Ser	Leu 355	Phe	Phe	Tyr	Pro	Ser 360	Tyr	Pro	Asp	Glu	Val 365	Arg	Gly	Val
Phe	Ala 370	Val	Arg	Thr	Gly	Lys 375	Tyr	Lys	Ala	His	Phe 380	Phe	Thr	Gln	Gly
Ser 385	Ala	His	Ser	Asp	Thr 390	Thr	Ala	Asp	Pro	Ala 395	Cys	His	Ala	Ser	Ser 400
Ser	Leu	Thr	Ala	His 405	Glu	Pro	Pro	Leu	Leu 410	Tyr	Asp	Leu	Ser	Lys 415	Asp
Pro	Gly	Glu	Asn 420	Tyr	Asn	Leu	Leu	Gly 425	Gly	Val	Ala	Gly	Ala 430	Thr	Pro
Glu	Val	Leu 435	Gln	Ala	Leu	Lys	Gln 440	Leu	Gln	Leu	Leu	Lys 445	Ala	Gln	Leu
Asp	Ala 450	Ala	Val	Thr	Phe	Gly 455	Pro	Ser	Gln	Val	Ala 460	Arg	Gly	Glu	Asp
Pro 465	Ala	Leu	Gln	Ile	Cys 470	Cys	His	Pro	Gly	Cys 475	Thr	Pro	Arg	Pro	Ala 480
Cys	Cys	His	Cys	Pro 485	Asp	Pro	His	Ala							